



ATTN: OFFICIAL DRAFTSPERSON
PATENT
Customer No. 22,852
Attorney Docket No. 08702.0005-00000

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

Lorenz et al.

Serial No.: 09/816,697

Filed: March 23, 2001

For: A NOVEL P-SELECTIN
GLYCOPROTEIN LIGAND
(PSGL-1) BINDING PROTEIN
AND USES THEREFOR

)
)
) **Notice of Draftsperson Dated:**
) **November 14, 2002**
)
) Group Art Unit: 1644
)
) Examiner: Phillip Gambel
)
)
)

ATTN: OFFICIAL DRAFTSPERSON
Director of the Patent and Trademark Office
Washington, DC 20231

Sir:

RECEIVED
APR 10 2003
TECH CENTER 1600/2900

SUBMISSION OF CORRECTED DRAWINGS

Please replace two sheet of drawings labeled as "FIGURE 1" and "FIGURE 1 cont'd" with the two (2) sheets of corrected drawings filed herewith (Fig. 1A and Fig. 1B). If the corrected drawings for any reason are not in full compliance with the pertinent statutes and regulations, please so advise the undersigned.

No fees are believed to be due for the submission of these drawings. If any fees are necessary, please charge Deposit Account No. 06-0916.

Respectfully submitted,

FINNEGAN, HENDERSON, FARABOW,
GARRETT & DUNNER, L.L.P.

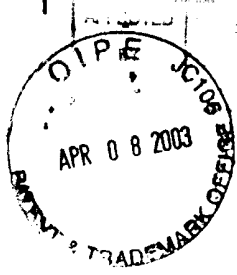
Dated: April 8, 2003

By: Leslie A. McDonell

Leslie A. McDonell
Reg. No. 34,872

FINNEGAN
HENDERSON
FARABOW
GARRETT &
DUNNER LLP

1300 I Street, NW
Washington, DC 20005
202.408.4000
Fax 202.408.4400
www.finnegan.com



RECEIVED
APR 10 2003
TECH CENTER 1600/2900

atg gca agt cca gag cac cct ggg agc cct ggc tgc atg gga cca gaa 48
Met Ala Ser Pro Glu His Pro Gly Ser Pro Gly Cys Met Gly Pro Ile
1 5 10 15

acc cag tgc acg gca agg acc cag cag gaa gca cca gcc act ggc ccc 96
Thr Gln Cys Thr Ala Arg Thr Gln Gln Glu Ala Pro Ala Thr Gly Pro
20 25 30

gac ctc ccg cac cca gga cct gac ggg cac tta gac aca cac agt ggc 144
Asp Leu Pro His Pro Gly Pro Asp Gly His Leu Asp Thr His Ser Gly
35 40 45

ctg agc tcc aac tcc agc atg acc acg cgg gag ctt cag cag tac tgg 192
Leu Ser Ser Asn Ser Ser Met Thr Thr Arg Glu Leu Gln Gln Tyr Trp
50 55 60

cag aac cag aaa tgc cgc tgg aag cac gtc aaa ctg ctc ttt gag atc 240
Gln Asn Gln Lys Cys Arg Trp Lys His Val Lys Leu Leu Phe Glu Ile
65 70 75 80

gct tca gct cgc atc gag gag aga aaa gtc tct aag ttt gtg gtg tac 288
Ala Ser Ala Arg Ile Glu Glu Arg Lys Val Ser Lys Phe Val Val Tyr
85 90 95

caa atc atc gtc atc cag act ggg agc ttt gac aac aac aag gcc gtc 336
Gln Ile Ile Val Ile Gln Thr Gly Ser Phe Asp Asn Asn Lys Ala Val
100 105 110

ctg gaa cgg cgc tat tcc gac ttc gcg aag ctc cag aaa gcg ctg ctg 384
Leu Glu Arg Arg Tyr Ser Asp Phe Ala Lys Leu Gln Lys Ala Leu Leu
115 120 125

aag acg ttc agg gag gag atc gaa gac gtg gag ttt ccc agg aag cac 432
Lys Thr Phe Arg Glu Glu Ile Glu Asp Val Glu Phe Pro Arg Lys His
130 135 140

ctg act ggg aac ttc gct gag gag atg atc tgt gag cgt cgg cgc gcc 480
Leu Thr Gly Asn Phe Ala Glu Glu Met Ile Cys Glu Arg Arg Arg Ala
145 150 155 160

ctg cag gag tac ctg ggc ctg ctc tac gcc atc cgc tgc gtg cgc cgc 528
Leu Gln Glu Tyr Leu Gly Leu Leu Tyr Ala Ile Arg Cys Val Arg Arg
165 170 175

tcc cgg gag ttc ctg gac ttc ctc acg cgg ccg gag ctg cgc gag gct 576
Ser Arg Glu Phe Leu Asp Phe Leu Thr Arg Pro Glu Leu Arg Glu Ala
180 185 190

ttc ggc tgc ctg cgg gcc ggc cag tac ccg cgc gcc ctg gag ctg ctg 624
Phe Gly Cys Leu Arg Ala Gly Gln Tyr Pro Arg Ala Leu Glu Leu Leu
195 200 205

ctg cgc gtg ctg ccg ctg cag gag aag ctc acc gcc cac tgc cct gcg 672
Leu Arg Val Leu Pro Leu Gln Glu Lys Leu Thr Ala His Cys Pro Ala
210 215 220

gcc gcc gtc ccg gcc ctg tgc gcc gtg ctg ctg tgc cac cgc gac ctc 720
Ala Ala Val Pro Ala Leu Cys Ala Val Leu Leu Cys His Arg Asp Leu
225 230 235 240

FIGURE 1A

gac cgc ccc gcc gag gcc ttc gcg gcc gga gag agg gcc ctg cag cgc	768
Asp Arg Pro Ala Glu Ala Phe Ala Ala Gly Glu Arg Ala Leu Gln Arg	
245 250 255	
ctg cag gcc cgg gag ggc cat cgc tac tat gcg cct ctg ctg gac gcc	816
Leu Gln Ala Arg Glu Gly His Arg Tyr Tyr Ala Pro Leu Leu Asp Ala	
260 265 270	
atg gtc cgc ctg gcc tac gcg ctg ggc aag gac ttc gtg act ctg cag	864
Met Val Arg Leu Ala Tyr Ala Leu Gly Lys Asp Phe Val Thr Leu Gln	
275 280 285	
gag agg ctg gag gag agc cag ctc cgg agg ccc acg ccc cga ggc atc	912
Glu Arg Leu Glu Glu Ser Gln Leu Arg Arg Pro Thr Pro Arg Gly Ile	
290 295 300	
acc ctg aag gag ctc act gtg cga gaa tac ctg cac tga	951
Thr Leu Lys Glu Leu Thr Val Arg Glu Tyr Leu His	
305 310 315	

FIGURE 1B